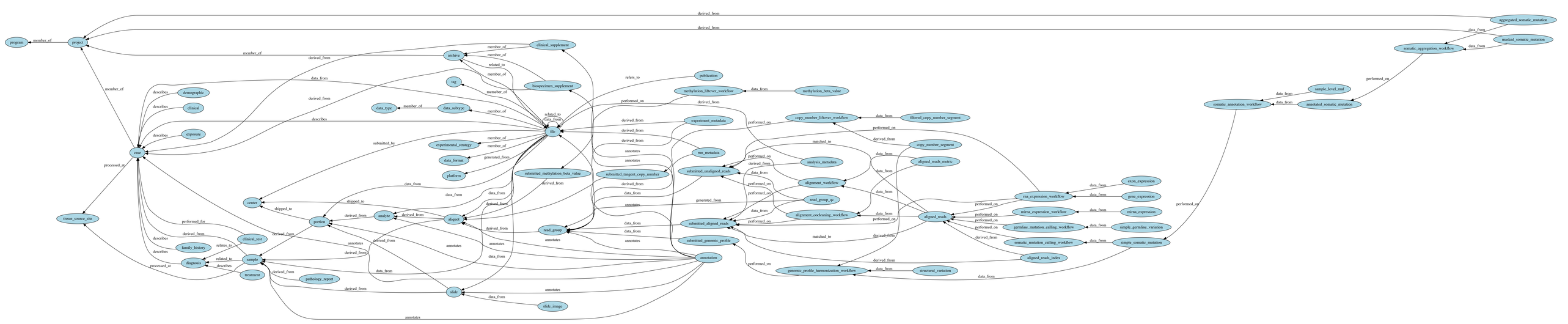
program
project
case
tissue_source_site
demographic
clinical
exposure
clinical_test
family_history
diagnosis
sample
treatment
pathology_report
center
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slide
slide_image
archive
tag
data_type
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experimental_strategy
data_format
platform
clinical_supplement
biospecimen_supplement
file
submitted_methylation_beta_value
submitted_tangent_copy_number
publication
experiment_metadata
run_metadata
analysis_metadata
read_group
read_group_qc
alignment_workflow
alignment_cocleaning_workflow
submitted_unaligned_reads
submitted_aligned_reads
submitted_genomic_profile
annotation
methylation_liftover_workflow
methylation_beta_value
copy_number_liftover_workflow
filtered_copy_number_segment
copy_number_segment
aligned_reads_metric
aligned_reads
rna_expression_workflow
exon_expression
gene_expression
mirna_expression_workflow
mirna_expression
germline_mutation_calling_workflow
simple_germline_variation
somatic_mutation_calling_workflow
simple_somatic_mutation
aligned_reads_index
genomic_profile_harmonization_workflow
structural_variation
aggregated_somatic_mutation
masked_somatic_mutation
somatic_aggregation_workflow
sample_level_maf
somatic_annotation_workflow
annotated_somatic_mutation
member_of
derived_from
describes
related_to
data_from
submitted_by
processed_at
processed_on
performed_for
relates_to
shipped_to
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refers_to
performed_on
annotates
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